



## ANNUAL OF PATENT EXAMINING PROCEDURE

PTO/SB/08 (2-92)  
Sheet 1 of 1

Form PTO-1449		Docket Number (Optional) 3087.00013		Application Number 10/780,318		
INFORMATION DISCLOSURE CITATION IN AN APPLICATION  (Use several sheets if necessary)				Applicant Hyesook Kim, et al.		
				Filing Date 02-17-04		
				Group Art Unit 1648		
U.S. PATENT DOCUMENTS						
EXAMINER INITIAL	DOCUMENT NUMBER	DATE	NAME	CLASS	SUBCLASS	FILING DATE IF APPROPRIATE
FOREIGN PATENT DOCUMENTS						
	DOCKET NUMBER	DATE	COUNTRY	CLASS	SUBCLASS	TRANSLATION YES NO
OTHER DOCUMENTS (Including Author, Title, Date Pertinent Pages, Etc.)						
SBC		Bantle, John A. et al. "Phase III Interlaboratory Study of FETAX Part 3. FETAX Validation using 12 Compounds with and without an Exogenous Metabolic Activation System," <i>Journal of Applied Toxicology</i> , 19:447-472 (1999).				
SBC		Fort, Douglas J. et al. "Evaluation of the Developmental Toxicity of Thalidomide Using Frog Embryo Teratogenesis Assay - Xenopus (FETAX): Biotransformation and Detoxification," <i>Teratogenesis, Carcinogenesis, and Mutagenesis</i> , 20:35-47 (2000).				
SBC		Newport, John et al. "Major Developmental Transition in Early Xenopus Embryos: I. Characterization and Timing of Cellular Changes at the Midblastula Stage," <i>Cell</i> , Vol. 30:675-686 (1982).				
SBC		Newport, John et al. "A Major Developmental Transition in Early Xenopus Embryos: II. Control of the Onset of Transcription," <i>Cell</i> , Vol. 30:657-696 (1982).				
SBC		Nakakura, Norihiko et al. "Synthesis of Heterogeneous mRNA-like RNA and Low-Molecular-Weight RNA before the Midblastula Transition in Embryos of <i>Xenopus laevis</i> ," <i>Developmental Biology</i> , 123:421-429 (1987).				
SBC		Adams, Mark D. et al. "Complementary DNA Sequencing: Expressed Sequence Tags and Human Genome Project," 1651-1656. (1991)				
SBC		Schena, Mark, et al. "Quantitative Monitoring of Gene Expression Patterns with a Complementary DNA Microarray," <i>Science</i> , Vol. 270:467-470 (1995).				
SBC		Schena, Mark et al. "Parallel human genome analysis: Microarray-based expression monitoring of 1000 genes," <i>Proc. Natl. Acad. Sci. USA</i> , Vol. 93:10614-10619 (1996).				
SBC		Zhang, Michael Q. "Large-Scale Gene Expression Data Analysis: A New Challenge to Computational Biologists," <i>Cold Spring Harbor Laboratory</i> , Vol. 9, Issue 8:681-688 (1999).				
SBC		Nuwaysir, Emile F. et al. "Microarrays and Toxicology: The Advent of Toxicogenomics," <i>Molecular Carcinogenesis</i> , 24:153-159 (1999).				
SBC		Altmann, Curtis R. et al. "Microarray-Based Analysis of Early Development in <i>Xenopus laevis</i> ," <i>Developmental Biology</i> , 236:64-75 (2001).				
SBC		External and Internal Stage Criteria in the Development of <i>Xenopus laevis</i> , Chapter VII, 162-188.				
		Nieuwkoop and Faber (1967)				
EXAMINER /Stacy B. Chen/ (11/30/2006)			DATE CONSIDERED 11/30/2006			
EXAMINER: Initial if citation considered, whether or not citation is in conformance with MPEP § 609. Draw line through citation if not in conformance and not considered. Include copy of this form with next communication to the applicant.						